WHAT IS CLAIMED IS:

1. A method for producing antimicrobial protein, comprising

expressing as a fusion protein in a prokaryotic cell, a basic antimicrobial protein A having a predetermined mode of an intramolecular disulfide bond as an active type by combining the protein A with a partner protein B having an isoelectric point below pH 7 and a chaperon function,

recovering the fusion protein, and

modifying and activating the antimicrobial protein A in the fusion protein into the active type by utilizing a function of the partner protein B.

- 2. The method according to claim 1, wherein the fusion protein is expressed by culturing a prokaryotic cell inserted with DNA encoding the fusion protein to express the DNA in the prokaryotic cell.
- 3. The method according to claim 1, wherein the antimicrobial protein A is any one of thionin, PR protein, lipid transfer protein and ribosome– inactivated protein, all derived from plants, or any one of diffensin derived from plants, insects and humans.
- 4. The method according to claim 1, wherein the partner protein B comprises an acid partner protein B1 at least with an isoelectric point below pH 7 and a chaperon partner protein B2 at least with a chaperon function.
- 5. The method according to claim 1, further comprising separating the antimicrobial protein A from the partner protein B in the fusion protein and modifying the antimicrobial protein A into an antimicrobially active type by utilizing the function of the partner protein B.
- 6. The method according to claim 5, wherein the step of separating

the antimicrobial protein A from the partner protein B includes the cleavage of a peptide bond in the border of the two proteins.

- 7. The method according to claim 5, wherein the step of separating the antimicrobial protein A from the partner protein B includes the cleavage of an oligopeptide moiety interposed in the border of the two proteins for cleavage.
- 8. The method according to claim 5, wherein the step of modifying the antimicrobial protein A into the antimicrobially active type thereof includes a general refolding procedure of disulfide bond for promoting the modification.
- 9. A fusion protein comprising

a basic antimicrobial protein A having a predetermined mode of an intramolecular disulfide bond as an active type, and

a partner protein B having an isoelectric point below pH 7 and a chaperon function.

- 10. The fusion protein according to claim 9, wherein the antimicrobial protein A and the partner protein B are chemically bonded together.
- 11. The fusion protein according to claim 9, wherein the antimicrobial protein A and the partner protein B form in series a polypeptide chain through an oligopeptide moiety enzymatically cleavable.
- 12. The fusion protein according to claim 9, wherein the antimicrobial protein A and the partner protein B are partially or wholly associated together via hydrophobic affinity or electric properties.
- 13. The fusion protein according to claim 9, wherein the antimicrobial protein A is any one of thionin, PR protein, lipid transfer protein and

ribosome-inactivating protein, all derived from plants, or any one of different derived from plants, insects and humans.

- 14. The fusion protein according to claim 9, wherein the partner protein B is thioredoxin (Tx) or chaperonin.
- 15. The fusion protein according to claim 9, wherein the chaperon function of the partner protein B is a refolding function to modify a wrong bonding position of the intramolecular disulfide bond into a right bonding position in the protein A for the active type thereof.
- 16. The fusion protein according to claim 9, wherein the partner protein B is protein disulfide isomerase (PDI) or an acid protein encoded by DNA downstream of the nucleotide sequence of thionin derived from plants.
- 17. A fusion protein comprising: a basic antimicrobial protein A having a predetermined mode of an intramolecular disulfide bond as an active type; an acid partner protein B1 at least with an isoelectric point below pH 7; and a chaperon partner protein B2 at least with a chaperon function.
- 18. The fusion protein according to claim 17, wherein the acid partner protein B1 comprises a carboxyl terminal region of the PDI derived from Furnicola insolens and the chaperon partner protein B2 is peptidylprolylcis-trans-isomerase.
- 19. A partner protein comprising an acid partner protein B1 at least with an isoelectric point below pH 7 and a chaperon partner protein B2 at least with a chaperon function, wherein the partner protein is a protein to be used for the formation of a fusion protein, together with a basic antimicrobial protein A having a predetermined mode of an intramolecular disulfide bond as an active type.

20. DNA encoding a fusion protein according to claim 1.